RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: _10/551,550_
Source: _/FwP_
Date Processed by STIC: _2/21/07_

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RAW SEQUENCE LISTING   DATE: 02/21/2007
PATENT APPLICATION: US/10/551,550   TIME: 14:20:03

Input Set : A:\10-551,550 Sequence Listing.txt
Output Set: N:\CRF4\02212007\J551550.raw

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 3 <110> APPLICANT: Seikagaku Corporation
 5 <120> TITLE OF INVENTION: Catalyst for cleaving sugar chain
 7 <130> FILE REFERENCE: Q90646
 9 <140> CURRENT APPLICATION NUMBER: US 10/551,550
10 <141> CURRENT FILING DATE: 2005-09-30
12 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/004695
13 <151> PRIOR FILING DATE: 2004-03-31
15 <150> PRIOR APPLICATION NUMBER: JP 2003-097301
16 <151> PRIOR FILING DATE: 2003-03-31
18 <150> PRIOR APPLICATION NUMBER: JP 2003-113965
19 <151> PRIOR FILING DATE: 2003-04-18
21 <160> NUMBER OF SEQ ID NOS: 2
22 <170> SOFTWARE: PatentIn version 3.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 1308
26 <212> TYPE: DNA
27 <213> ORGANISM: Homo sapiens
29 <220> FEATURE:
30 <221> NAME/KEY: CDS
31 <222> LOCATION: (1)..(1308)
33 <400> SEQUENCE: 1
34 atg gca gcc cac ctg ctt ccc atc tgc gcc ctc ttc ctg acc tta ctc      48
35 Met Ala Ala His Leu Leu Pro Ile Cys Ala Leu Phe Leu Thr Leu Leu
36 1               5                   10                  15
37 gat atg gcc caa ggc ttt agg ggc ccc ttg cta ccc aac cgg ccc ttc      96
38 Asp Met Ala Gln Gly Phe Arg Gly Pro Leu Leu Pro Asn Arg Pro Phe
39                 20                  25                  30
40 acc acc gtc tgg aat gca aac acc cag tgg tgc ctg gag agg cac ggt     144
41 Thr Thr Val Trp Asn Ala Asn Thr Gln Trp Cys Leu Glu Arg His Gly
42         35                  40                  45
43 gtg gac gtg gat gtc agt gtc ttc gat gtg gta gcc aac cca ggg cag     192
44 Val Asp Val Asp Val Ser Val Phe Asp Val Val Ala Asn Pro Gly Gln
45         50                  55                  60
46 acc ttc cgc ggc cct gac atg aca att ttc tat agc tcc cag ctg ggc     240
47 Thr Phe Arg Gly Pro Asp Met Thr Ile Phe Tyr Ser Ser Gln Leu Gly
48 65                  70                  75                  80
49 acc tac ccc tac tac acg ccc act ggg gag cct gtg ttt ggt ggt ctg     288
50 Thr Tyr Pro Tyr Tyr Thr Pro Thr Gly Glu Pro Val Phe Gly Gly Leu
51                 85                  90                  95
52 ccc cag aat gcc agc ctg att gcc cac ctg gcc cgc aca ttc cag gac     336
53 Pro Gln Asn Ala Ser Leu Ile Ala His Leu Ala Arg Thr Phe Gln Asp
54                 100                 105                 110
55 atc ctg gct gcc ata cct gct cct gac ttc tca ggg ctg gca gtc atc     384
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56 Ile Leu Ala Ala Ile Pro Ala Pro Asp Phe Ser Gly Leu Ala Val Ile
57         115             120             125
58 gac tgg gag gca tgg cgc cca cgc tgg gcc ttc aac tgg gac acc aag      432
59 Asp Trp Glu Ala Trp Arg Pro Arg Trp Ala Phe Asn Trp Asp Thr Lys
60     130             135             140
61 gac att tac cgg cag cgc tca cgg gca ctg gta cag gca cag cac cct      480
62 Asp Ile Tyr Arg Gln Arg Ser Arg Ala Leu Val Gln Ala Gln His Pro
63 145             150             155             160
64 gat tgg cca gct cct cag gtg gag gca gta gcc cag gac cag ttc cag      528
65 Asp Trp Pro Ala Pro Gln Val Glu Ala Val Ala Gln Asp Gln Phe Gln
66             165             170             175
67 gga gct gca cgg gcc tgg atg gca ggc acc ctc cag ctg ggg cgg gca      576
68 Gly Ala Ala Arg Ala Trp Met Ala Gly Thr Leu Gln Leu Gly Arg Ala
69         180             185             190
70 ctg cgt cct cgc ggc ctc tgg ggc ttc tat ggc ttc cct gac tgc tac      624
71 Leu Arg Pro Arg Gly Leu Trp Gly Phe Tyr Gly Phe Pro Asp Cys Tyr
72         195             200             205
73 aac tat gac ttt cta agc ccc aac tac acc ggc cag tgc cca tca ggc      672
74 Asn Tyr Asp Phe Leu Ser Pro Asn Tyr Thr Gly Gln Cys Pro Ser Gly
75         210             215             220
76 atc cgt gcc caa aat gac cag cta ggg tgg ctg tgg ggc cag agc cgt      720
77 Ile Arg Ala Gln Asn Asp Gln Leu Gly Trp Leu Trp Gly Gln Ser Arg
78 225             230             235             240
79 gcc ctc tat ccc agc atc tac atg ccc gca gtg ctg gag ggc aca ggg      768
80 Ala Leu Tyr Pro Ser Ile Tyr Met Pro Ala Val Leu Glu Gly Thr Gly
81             245             250             255
82 aag tca cag atg tat gtg caa cac cgt gtg gcc gag gca ttc cgt gtg      816
83 Lys Ser Gln Met Tyr Val Gln His Arg Val Ala Glu Ala Phe Arg Val
84             260             265             270
85 gct gtg gct gct ggt gac ccc aat ctg ccg gtg ctg ccc tat gtc cag      864
86 Ala Val Ala Ala Gly Asp Pro Asn Leu Pro Val Leu Pro Tyr Val Gln
87         275             280             285
88 atc ttc tat gac acg aca aac cac ttt ctg ccc ctg gat gag ctg gag      912
89 Ile Phe Tyr Asp Thr Thr Asn His Phe Leu Pro Leu Asp Glu Leu Glu
90     290             295             300
91 cac agc ctg ggg gag agt gcg gcc cag ggg gca gct gga gtg gtg ctc      960
92 His Ser Leu Gly Glu Ser Ala Ala Gln Gly Ala Ala Gly Val Val Leu
93 305             310             315             320
94 tgg gtg agc tgg gaa aat aca aga acc aag gaa tca tgt cag gcc atc     1008
95 Trp Val Ser Trp Glu Asn Thr Arg Thr Lys Glu Ser Cys Gln Ala Ile
96             325             330             335
97 aag gag tat atg gac act aca ctg ggg ccc ttc atc ctg aac gtg acc     1056
98 Lys Glu Tyr Met Asp Thr Thr Leu Gly Pro Phe Ile Leu Asn Val Thr
99             340             345             350
100 agt ggg gcc ctt ctc tgc agt caa gcc ctg tgc tcc ggc cat ggc cgc    1104
101 Ser Gly Ala Leu Leu Cys Ser Gln Ala Leu Cys Ser Gly His Gly Arg
102         355             360             365
103 tgt gtc cgc cgc acc agc cac ccc aaa gcc ctc ctc ctc ctt aac cct    1152
104 Cys Val Arg Arg Thr Ser His Pro Lys Ala Leu Leu Leu Leu Asn Pro
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Input Set : A:\10-551,550 Sequence Listing.txt
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105     370                 375                 380
106 gcc agt ttc tcc atc cag ctc acg cct ggt ggt ggg ccc ctg agc ctg    1200
107 Ala Ser Phe Ser Ile Gln Leu Thr Pro Gly Gly Gly Pro Leu Ser Leu
108 385                 390                 395                 400
109 cgg ggt gcc ctc tca ctt gaa gat cag gca cag atg gct gtg gag ttc    1248
110 Arg Gly Ala Leu Ser Leu Glu Asp Gln Ala Gln Met Ala Val Glu Phe
111                 405                 410                 415
112 aaa tgt cga tgc tac cct ggc tgg cag gca ccg tgg tgt gag cgg aag    1296
113 Lys Cys Arg Cys Tyr Pro Gly Trp Gln Ala Pro Trp Cys Glu Arg Lys
114                 420                 425                 430
115 agc atg tgg tga                                                    1308
116 Ser Met Trp
117         435
120 <210> SEQ ID NO: 2
121 <211> LENGTH: 435
122 <212> TYPE: PRT
123 <213> ORGANISM: Homo sapiens
125 <400> SEQUENCE: 2
126 Met Ala Ala His Leu Leu Pro Ile Cys Ala Leu Phe Leu Thr Leu Leu
127 1               5                   10                  15
128 Asp Met Ala Gln Gly Phe Arg Gly Pro Leu Leu Pro Asn Arg Pro Phe
129             20                  25                  30
130 Thr Thr Val Trp Asn Ala Asn Thr Gln Trp Cys Leu Glu Arg His Gly
131         35                  40                  45
132 Val Asp Val Asp Val Ser Val Phe Asp Val Val Ala Asn Pro Gly Gln
133     50                  55                  60
134 Thr Phe Arg Gly Pro Asp Met Thr Ile Phe Tyr Ser Ser Gln Leu Gly
135 65                  70                  75                  80
136 Thr Tyr Pro Tyr Tyr Thr Pro Thr Gly Glu Pro Val Phe Gly Gly Leu
137                 85                  90                  95
138 Pro Gln Asn Ala Ser Leu Ile Ala His Leu Ala Arg Thr Phe Gln Asp
139             100                 105                 110
140 Ile Leu Ala Ala Ile Pro Ala Pro Asp Phe Ser Gly Leu Ala Val Ile
141         115                 120                 125
142 Asp Trp Glu Ala Trp Arg Pro Arg Trp Ala Phe Asn Trp Asp Thr Lys
143     130                 135                 140
144 Asp Ile Tyr Arg Gln Arg Ser Arg Ala Leu Val Gln Ala Gln His Pro
145 145                 150                 155                 160
146 Asp Trp Pro Ala Pro Gln Val Glu Ala Val Ala Gln Asp Gln Phe Gln
147                 165                 170                 175
148 Gly Ala Ala Arg Ala Trp Met Ala Gly Thr Leu Gln Leu Gly Arg Ala
149             180                 185                 190
150 Leu Arg Pro Arg Gly Leu Trp Gly Phe Tyr Gly Phe Pro Asp Cys Tyr
151         195                 200                 205
152 Asn Tyr Asp Phe Leu Ser Pro Asn Tyr Thr Gly Gln Cys Pro Ser Gly
153     210                 215                 220
154 Ile Arg Ala Gln Asn Asp Gln Leu Gly Trp Leu Trp Gly Gln Ser Arg
155 225                 230                 235                 240
156 Ala Leu Tyr Pro Ser Ile Tyr Met Pro Ala Val Leu Glu Gly Thr Gly
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```
157                 245                 250                 255
158 Lys Ser Gln Met Tyr Val Gln His Arg Val Ala Glu Ala Phe Arg Val
159                 260                 265                 270
160 Ala Val Ala Ala Gly Asp Pro Asn Leu Pro Val Leu Pro Tyr Val Gln
161         275                 280                 285
162 Ile Phe Tyr Asp Thr Thr Asn His Phe Leu Pro Leu Asp Glu Leu Glu
163         290                 295                 300
164 His Ser Leu Gly Glu Ser Ala Ala Gln Gly Ala Ala Gly Val Val Leu
165 305                 310                 315                 320
166 Trp Val Ser Trp Glu Asn Thr Arg Thr Lys Glu Ser Cys Gln Ala Ile
167                 325                 330                 335
168 Lys Glu Tyr Met Asp Thr Thr Leu Gly Pro Phe Ile Leu Asn Val Thr
169                 340                 345                 350
170 Ser Gly Ala Leu Leu Cys Ser Gln Ala Leu Cys Ser Gly His Gly Arg
171         355                 360                 365
172 Cys Val Arg Arg Thr Ser His Pro Lys Ala Leu Leu Leu Leu Asn Pro
173         370                 375                 380
174 Ala Ser Phe Ser Ile Gln Leu Thr Pro Gly Gly Gly Pro Leu Ser Leu
175 385                 390                 395                 400
176 Arg Gly Ala Leu Ser Leu Glu Asp Gln Ala Gln Met Ala Val Glu Phe
177                 405                 410                 415
178 Lys Cys Arg Cys Tyr Pro Gly Trp Gln Ala Pro Trp Cys Glu Arg Lys
179                 420                 425                 430
180 Ser Met Trp
181         435
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VERIFICATION SUMMARY   DATE: 02/21/2007
PATENT APPLICATION: **US/10/551,550**   TIME: 14:20:04

Input Set : **A:\10-551,550 Sequence Listing.txt**
Output Set: **N:\CRF4\02212007\J551550.raw**